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Database :
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Maximum DB seq length: 2000000000
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Maximum Match 100%
Listing first 45 summaries
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1937.259 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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## ALIGNMENTS

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(BALT/) BALTIMORE D. (CELEN/) CHENG G. (YEZZ/) YE Z. (LEDE/) LEDERMAN S. (CLEA/) CLEARY A.		14-MAR-2002. 10-MAR-1997; 97US-0813323.	US2002031522-A1.	antipsoriatic; bladder disease; human herpesvirus 4; Epstein-Barr virus. Mus sp.	antianaemic; anti-thyroid; anti-heumatic; antiarthritic; cardiant; dermatological; haemostatic; antidiabetic; antiarteriosclerotic;	autoimmune disease; paptocare disease; pastrointestinal disease; vasotropic; finance disease; vasotropic;		15-AUG-2002 (first entry)	AA017756;	LT 1 7756 AAO17756 standard; protein; 567 AA.

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YDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQT
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Best Local Similarity
                                                                                                                                               Matches
                                                                                                                                                                                                                                                conditions, comprising measuring the expression levels of particular genes (I) in a test sample or determining the expression profile of a gene group in the sample comprising genes selected from (I). The method is useful for examining the ischaemia genes selected from (I). The method is useful for examining the ischaemia or vasospastic ischaemia) by measuring expression levels of particular genes (AB199202 to AB199012, encoding the protein sequences in ABB57020 to ABB57374) or by determining the expression profile of a gene group comprising these genes. The expression levels or expression profiles produced by these genes are used as an indicator when screening for ischaemic condition improving drugs or therapeutics for ischaemic diseases. AB199913 and AB19914 represent PCR primers for a mouse ischaemic condition related sequence, which are used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABB57054 standard; Protein; 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Examining the ischemic condition (e.g. occlusive ischemia) by measuring expression levels of particular genes defined in the specification or by determining the expression profile of a gene group comprising these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB57054;
                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a method for examining ischaemic conditions, comprising measuring the expression levels of particular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 297-300; 2690pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; ABI99264.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-MAY-2001; 2001WO-JP04192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200188188-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vasospastic ischaemia; ischaemic condition; ischaemic disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse; ischaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse ischaemic condition related protein sequence SEQ ID NO:98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genes
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                                                                                         1 MESSKKMDAAGTLQPNPPLKLQPDRGAGSVLVPEQGGYKEKFVKTVEDKYKCEKCRLVLC 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NPKQTECGHRFWQSCMAALLSSSSPKCTACQESIIKDKVFKDNCCKREILALQVYCRNEG
                                NPKQTECGHRFCESCMAALLSSSSPKCTACQESIIKDKVFKDNCCKREILALQVYCRNEG 120
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                                                                    \tt MESSKKMDAAGTLQPNPPLKLQPDRGAGSVLVPEQGGYKEKFVKTVEDKYKCEKCRLVLC
                                                                                                                                               Conservative
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                                                                                                                                                              99.3%;
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                                                                                                                                                              Score 2974; DB 23;
Pred. No. 2.3e-239;
                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                      KW CD40 receptor associated factor 1; CRAF1-a; TRAF-3; p55; human; KW CD40 mediated intracellular signalling; organ rejection; allergy; KW hay fever; autoimmune disease; systemic lupus erythematosus; KW rheumatoid arthritis; myasthenia gravis; Graves' disease; KW idiopathic thrombocytopaenia purpura; haemolytic anaemia; KW diabetes mellitus; psoriasis; hyper immunoglobulin E syndrome; KW apoptosis; Reter's syndrome; spondyloarthritis; Lymon disease; HIV; KW apoptosis; steter's syndrome; spondyloarthritis; pummonary fibrosis; KW pneumoconiosis; arthritis; scleroderma; pulmonary fibrosis; KW apoptosis; siliconosis; arthritis; scleroderma; pulmonary fibrosis; KW abbestosis; siliconosis; farmer's lung; hepatitis; cirrhosis; KW atherosclerosis; glomerulopathy; kidney disease; nephropathy; KW endocarditis; leprosy; malaria; Goodpasture's disease; KW Henoch-Schoenlein purpura; polyarteritis; multiple myeloma; KW wegener's macroglobulinaemia; amyloidosis; Sjogren's sydrome; KW Maldenstroem's macroglobulinaemia; significantis; B cell tumour; KW Maldenstroem's macroglobulinaemia; spondader disease; Epstein-Barr virus; mononucleosis; B cell tumour; KW Multiphemalia; macronary rirus; mononucleosis; B cell tumour;
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                                                                                                                                                                                                                                                                                                                                                   Burkitt's lymphoma; nasopharyngeal carcinoma; pneumonia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human CRAF1-a (TRAF-3-p55) polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW27431 standard; Protein; 568
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                                                                                                                                                     Location/Qualifiers
117..141
                                                                   "zinc finger 1 (2n binding to Cys-117, Cys-124, His-136 and Cys-141)"
"zinc finger 2 (zinc binding to Cys-148
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61

60 CNPKQTECGHRFCESCMAALLSSSSPKCTACQESIIKDKVFKDNCCKREILALQVYCRNE 119

CSPKQTECGHRFCESCMAALLSSSSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNE 120

1 MESSKKMDAAGTLQPNPPLKLQPDRGAGS-VLVPEQGGYKEKFVKTVEDKYKCEKCRLVL 59 

밁

Matches Query Match Best Local

Similarity

96.4**%**; 96.1**%**;

Score 2886.5; DB 18; Length 568; Pred. No. 4.7e-232; 7; Mismatches 14; Indels 1;

1; Gaps

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Conservative

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CRAFI-a or TRAF-3-p55, p55, CRAFI(p55), TRAF-3(p55) or CRAFI(p60).
CC RAFI-a or TRAF-3-p55, p55, CRAFI(p55), TRAF-3(p55) or CRAFI(p60).
CC It is encoded by exons 4-13 of the human CRAF gene (see AAT90123).
CC CRAFI-a is a signalling protein that interacts with the cytoplasmic CC tail of B cell surface molecule CD40 and mediates a variety of T-dependent effects on B cell activation and differentiation. A CC higher mol.wt. CRAFI, designated CRAFID (see AAW27428), has also cell activation of Zinc Fingers. CRAFID that comprise different combinations of Zinc fingers. CRAFI epetides, CC comprise different combinations of Zinc fingers. CRAFI epetides, CC comprise different combinations of Zinc fingers. CRAFI epetides, CC comprise different combinations of Zinc fingers. CRAFI epetides, CC comprise different combinations of Zinc fingers. CRAFI epetides, CC comprise different combinations of Zinc fingers. CRAFI epetides, CC comprise different on their surface, particularly by introducing the CC characterised by an aberrant or unwanted level of CD40 mediated CC intracellular signalling, such as organ rejection, or a CD40 condition may be an allergic response or an autoimmune response, or CC condition may be an allergic response or an autoimmune response, or cells, an inflammatory kidney disease, a smooth muscle cell-dependent disease, or a condition associated with Epstein-Barr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-SEP-1996;
21-MAR-1996;
01-MAY-1996;
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protein comprising CRAFI-b domain capable of inhibiting CD40 mediated cell activation - useful to treat conditions characterised
                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Fig 1D-O; 158pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cleary AM, Frank DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               by aberrant or unwanted level of CD40 mediated intracellular
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568 AA;
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96US-0016626
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "zinc finger 4 (2n binding to Cys-208, His-221 and Cys-225)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "zinc finger 5 (2n binding to Cys-232, Cys-239, His-251 and 259-381"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cys-153, His-165 and Cys-170'
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RESULT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide; phosphorothioate; antiproliferative; anti-inflammatory; E-selectin; jun kinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY98166 standard; Protein; 568 AA
                                                                                                                                                                                     Antisense oligonucleotides targeted to nucleic acids encoding human tumour necrosis factor receptor-associated factor (TRAF), useful for treating diseases associated with TRAF expression such as inflammatory
                                                                                                                                                                                                                                                                                                                      WPI; 2000-303732/26.
N-PSDB; AAA55492.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumour necrosis factor receptor-associated factor; TRAF; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human TRAF3 protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-AUG-2000
      The present invention relates to (see AAA55496-A55757) which are
                                                                                            Disclosure; Page 111-114; 170pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-OCT-1998;
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         targeted
         antisense oligonucleotides
targeted to nucleic acids encoding
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RESULT 5 AAO17757

AAO17757 standard; protein;

AAO17757; 15-AUG-2002

(first entry)

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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating a human having a disease associated with the expression of TRAF comprising administering an antisense oligonucleotide. The reduction of jun kinase activation in cells comprises contacting the cells with an antisense oligonucleotide targeted to TRAF-6. A method for the reduction of E-selectin expression in cells or tissues comprises contacting the cells or tissues with an antisense oligonucleotide targeted to TRAF-2 or TRAF-6. The antisense oligonucleotides have antiproliferative and anti-inflammatory activity and are useful for treating disorders associated with cell proliferation and inflammation. The antisense oligonucleotides may also be used as a diagnostic probe for studying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human tumour necrosis factor receptor-associated factor (TRAF). The antisense sequences comprise at least one modified internucleotide linkage, which is a phosphorothioate linkage. The oligonucleotides also include at least one modified sugar molety such as a 2'-O-methoxyethyl sugar molety. Sequences AAA55490-A55495 represent nucleotide sequences encoding human TRAF1-6, and sequences AAY98169 represent the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ocal Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR 359
                                                                                                                                                                                                                                                                                                                                                                                     FQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              KSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCV
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                                                                                                                                                    KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
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                                  TVLENGTYIKDDTIFIKVIVDTSDLPDP 567
                                                                                                           EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ
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TVLENGTYIKDDTIFIKVIVDTSDLPDP
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96.1%;
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                                                                                                                                                                                                                                                                                                                                          CC CD40-mediated intracellular signaling, such as: organ rejection, continuum e diseases such as rheumatoid arthritis, myasthaenia gravis, cc systemic lupus erythematosus, Grave's disease, idiopathic continuum e diseases such as rheumatoid arthritis, myasthaenia gravis, cc systemic lupus erythematosus, Grave's disease, idiopathic continuum e disease, activation of fibroblast cells (e.g. allergic response (e.g. hay fever or a penicillin allergy), a condition continuum e disease, a condition dependent on CD40. Cc ligand-induced activation of endothelial cells (e.g. atherosclerosis, cc reperfusion injury, allograft rejection, organ rejection, or chronic confilammatory autoimmune diseases, a condition dependent on CD40. Cc (e.g. paoriasis), or an inflammatory kidney disease (e.g. membranous condition dependent on cD40. Cc (e.g. paoriasis), or an inflammatory kidney disease (e.g. membranous condition dependent on cD40. Cc (e.g. paoriasis), or an inflammatory kidney disease (e.g. membranous condition dependent on cD40. Cc (e.g. paoriasis), or an inflammatory kidney disease (e.g. membranous condition dependent on cD40. Cc (e.g. paoriasis), or an inflammatory kidney disease (e.g. membranous condition dependent on cD40. Cc (e.g. paoriasis), or an inflammatory kidney disease (e.g. membranous condition dependent on cD40. Cc (e.g. paoriasis), or an inflammatory kidney disease (e.g. membranous condition dependent on cD40. Cc (e.g. paoriasis), or an inflammatory kidney disease (e.g. membranous cc (e.g. membranous c
                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a protein comprising a CD40 receptor-associated factor 1 (CRAF1) truncated by about 323 - 414 amino acid residues at the amino terminus, or its variant, which is capable of inhibiting CD40-mediated cell activation. The protein is useful for treating a condition characterised by an aberrant or unwanted level of condition characterised by an aberrant or unwanted level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; CD40 receptor-associated factor 1; CRAF1; organ rejection; autoimmune disease; apoptosis; infection; fibrosis; liver disease; kidney disease; vascular disease; gastrointestinal disease; vasotropic; immunosuppressive; antiinflammatory; nephtropic; antiallergic; antianaemic; anti-thyroid; antirheumatic; antiarthritic; cardiant; dermatological; haemostatic; antidiabetic; antiarteriosclerotic; antipsoriatic; bladder disease; human herpesvirus 4; Epstein-Barr virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New CD40 receptor-associated factor 1 capable of inhibiting CD40-mediated cell activation, useful for treating e.g. inflammatory diseases, autoimmune diseases, allergic reaction, or organ transplant
                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 1; 31pp; English.
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(W CD40 mediated intracellular signaling; organ rejection; allergy;

(W hay fever; autoimmune disease; systemic lupus erythematosus;

(W rheumatoid arthritis; myasthenia gravis; Graves' disease;

(W alopathic thrombocytopaenia purpura; haemolytic anaemia;

(W diabetes mellitus; psoriasis; hyper immunoglobulin E syndrome;

(W apoptosis; Rieter's syndrome; spondyloarthritis; Lyme disease; Hiv;

(W apoptosis; rieter's syndrome; spondyloarthritis; Lyme disease; Hiv;

(W apoptosis; adult respiratory distress syndrome; pneumonitis;

(W appellis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis;

(W appellis; tuberculosis; Farmer's lung; hepatitis; cirrhosis;

(W asbestosis; siliconosis; Farmer's lung; hepatitis; cirrhosis;

(W alomerulosclerosis; glomerulopathy; kidney disease; nephropathy;

(W alomerulosclerosis; malaria; Goodpasture's disease;

(W endocarditis; leprosy; malaria; Goodpasture's disease;

(W wegener's granulomatosis; cryoimmunoglobulinaemia;

(W wegener's granulomatosis; cryoimmunoglobulinaemia;

(W waldenstroem's macroglobulinaemia; amyloidosis; Sjogren's sydrome;

(W walder disease; Epstein-Barr virus; mononucleosis; B cell tumour;

(W wanoa thorany, diamonai; nasopharyngeal carcinoma; pneumonia;
                                                                                                                                                                                                                                            Homo sapiens.
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Region

/note=

"zinc finger 1 (Zn binding to Cys-239,

242

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Qy
                                                                                                                                                     This polypeptide comprises a CRAFI (TRAF-3) protein designated CC CRAFI-b or TRAF-3-p70 or p70 or CRAFI(p70) or TRAF-3(p70). It is encoded by exons 1-2 and 4-13 of the human CRAF gene (see a AF90123). A lower mol.wt. CRAFI, designated CRAFIa (see AAW27431), has calso been identified, as well as isoforms p5 (see AAW27429), p15 (see AAW27430) and variants of CRAFI as isoforms p5 (see AAW27429), p15 (see CAAW27430) and variants of CRAFI-fine and CRAF- (see AAW27429), p15 (see CAAW27430) and variants of CRAFI-fine and CRAF- (see AAW27429), p15 (see COMPISE (see CAAW27430)) and variants of CRAFI-fine see AAW27430-37) that comprise different combinations of zinc fingers. CRAFI peptides, comprise of the seed to inhibit CD40 ligand activation of cells that cexpress CD40 on their surface, particularly by introducing the nucleic acid molecule into the cells, useful to treat conditions contacted acid molecule into the cells, useful to treat conditions contacted acid molecule into the cells, useful to treat conditions contacted intracellular signalling, such as organ rejection, or a CD40 intracellular signalling, such as organ rejection, or a CD40 condition may be an allergic response or an autoimmune response, or may be dependent on CD40 ligand-induced activation of epithelial conditions and inflammatory kidney disease, a smooth muscle cell-
                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-SEP-1996;
21-MAR-1996;
01-MAY-1996;
01-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein comprising CRAFI-b domain capable of inhibiting CD40 mediated cell activation - useful to treat conditions characterised by aberrant or unwanted level of CD40 mediated intracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cleary AM,
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                                                                                                                                                     Sequence
                                                                                                                                                                                                                    dependent disease, or a condition associated with Epstein-Barr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Fig 1A-O; 158pp; English.
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                                                                 sal Similarity
546; Conser
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                                                                 Conservative
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96US-0016626.
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                                                                                   96.4%;
96.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lederman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "zinc finger 4 (Zn binding to Cys-326, 330, His-343 and Cys-347)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "putative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "zinc finger 3 (Zn binding to Cys-299, Cys-302, His-314 and Cys-319"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "zinc finger 2 (zinc binding to Cys-270,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "zinc finger 5 (Zn binding to Cys-354, Cys-361, His-373 and His-381"
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                                                                 7;
                                                          Score 2886.5; DB 10.
Pred. No. 6.2e-232;
Pred. No. 6.2e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            s
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AAW03146
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                                                                                                                                                                                                                                                                                                                                                                                                                tumour necrosis factor receptor associated factor; TRAF; signal transduction; TNF; TNFR; lymphoblast; tumorigenesis; AIDS; Hodgkin's disease; Burkitt's lymphoma; naspharyngeal carcinoma; mononnucleosis; Epstein-Barr virus; EBV; therapy.
                        30-DEC-1994;
                                                                                                                                                       W09620723-A1
                                                                                                                                                                                                                       Domain
                                                                                                                                                                                                                                                                  Domain
                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LAP1; LMP1 associated protein 1; latent infection membrane protein;
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                                                                   28-DEC-1995;
                                                                                                                                                                                                                                                                                                             Domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKYREATCSHC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLRFQVLETASYNGVLIW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHC
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                           94US-0367540
                                                                   95WO-US16980
                                                                                                                                                                                                                                                                  309..341
                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                           /Label= Coiled_coil_domain
                                                                                                                                                                                                                                                                                      /label= LMP1-Binding_domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A novel human B-cell protein (AAW03146), termed LMP1 associated protein 1 or LAP1, strongly associates with the cytoplasmic C-terminal domain (AAW03148) of Epstein-Barr virus (EBV) latent infection membrane protein 1 (LMP1), a domain that is stringently required for transformed cell growth. LAP1 is related to murine tumour necrosis factor receptor associated factor TRAF2. A related novel B-cell protein (AAW03147), EBIT6, appears to be the human homologue of murine TRAF1. LAP1 polypeptides, esp. the LMP1 binding domain, coiled coil domain and C-terminal domain can be used to inhibit LMP1-TRAF interaction. Such polypeptides, which may be obtd. by recombinant means (see also AAT31273) can be used to treat infection and control cell growth or tumourigenesis associated with LMP1-encoding viruses, partic. EBV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compounds and methods for controlling TRAF-mediated signals - by modulating interactions between Epstein Barr virus encoded proteins LMP1, TAP1, TNFR to inhibit lymphoblast growth and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 74; Page 58-60; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1996-333765/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Birkenbach M,
                   540
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TVLENGTYIKDDTIFIKVIVDTSDLPDP
                                                                                                                                   KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
                                                                                                                                                                                    VTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLREQVLETASYNGVLIW
                                                                                                                                                                                                                                                                    EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR
                                                                                                                                                                                                                                                                                                                                        FQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI
                                                                                                                                                                                                                                                                                                                                                                                                           KSQVPMIKLQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CSPKQTECGHRFCESCMAALLSSSSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CNPKQTECGHRFCESCMAALLSSSSPKCTACQESIIKDKVFKDNCCKREILALQVYCRNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MESSKKMDSPGALQTNPPLKLHTDRSAGTPVFVPEQGGYKEKFVKTVEDKYKCEKCHLVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MESSKKMDAAGTLQPNPPLKLQPDRGAGS-VLVPEQGGYKEKFVKTVEDKYKCEKCRLVL
                                                                  EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ
                                                                                                                   KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
                                                                                                                                                                                                                       VTELESVDKSAGQAARNTGLLESQLSRHDQTLSVHDIRLADMDLRFQVLETASYNGVLIW
                                                                                                                                                                                                                                                                                                                                                                                         KSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SRGCAEQLMLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKYREATCSHC
                                                  EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ
                                                                                                                                                                                                                                                    EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR
                                                                                                                                                                                                                                                                                                                         FQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                545;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BRIGHAM & WOMENS HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UNIV CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               568 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kaye KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96.2%;
96.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kieff E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 2880.5; DB 17;
Pred. No. 1.5e-231;
7; Mismatches 15; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mosialos G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vanarsdale
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 568;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       +
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                                                  54 C
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                                                                                                                   48C
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                                                                                                                                                                                                                                                                                                                         30C
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                                                                                                                                                                                                                                                      36C
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GRGCAEQLTLGHLLVHLKNECOFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHC 179

CSPKQTECGHRFCESCMAALLSSSSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNE

120

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SRGCAEQLMLGH-LVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKYREATCSHC

61

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RESULT 8
AAB67615
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                                                                                                                                Query Match
Best Local Similarity
                                                                                                                   Matches
                                                                                                                                                                                                          The present sequence represents a human CD40 ligand. CD40 ligands are used for reducing osteoblast cell death or apoptosis, and for treating or preventing bone loss in animals, preferably humans, at risk of, or undergoing, bone loss. The bone loss is associated with osteoporosis, osteonecrosis, inflammatory arthritis, post-menopausal oestrogen loss, esterogen loss due to ovariectomy, total historectomy, lupus nephritis, rakayasu's arteritis, wegener's granulomatosis, anti-glomerular basement thrombocytopenia, asthma, a chronic obstructive lung disease, nephrotic/nephritic syndrome, or cancer. They may also be used to treat or prevent bone loss in a subject undergoing, or scheduled for, an organ or bone marrow transplant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CD40 ligand; osteoblast cell death; apoptosis; bone loss; osteoporosis; osteonecrosis; inflammatory arthritis; esterogen lo: ovariectomy; historectomy; lupus nephritis; Takayasu's arteritis; Wegener's granulomatosis; nephritis; myositis; scleroderma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB67615 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 116-118; 118pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CD40 agonist containing composition, used to reduce bone cell death or apoptosis associated with osteoporosis, osteonecrosis and inflammatory arthritis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; AAF55540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ahuja SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200116180-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Amino acid sequence of human CD40 ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB67615;
                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-AUG-2000; 2000WO-US23276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thrombocytopenia; asthma; lung disease; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   541 TVLENGTYIKDDTIFIKVIVDTSDLPDP 568
                 60
                                                                                1 MESSKKMDAAGTLQPNPPLKLQPDRGAGS-VLVPEQGGYKEKFVKTVEDKYKCEKCRLVL 59
CNPKQTECGHRFCESCMAALLSSSSPKCTACQESIIKDKVFKDNCCKREILALQVYCRNE 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001-169007/17.
                                                 MESSKKMDSPGALQTNPPLKLHTDRSAGTPVFVPEQGGYKEKFVKTVEDKYKCEKCHLVL
                                                                                                                   543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bonewald LF;
                                                                                                                                                                                  567 AA;
                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0151250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protein;
                                                                                                                               95.5%;
95.6%;
                                                                                                               7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    567
                                                                                                                               Score 2859; DB 22; Pred. No. 9.2e-230;
                                                                                                                   Mismatches
                                                                                                                   Indels
                                                                                                                                                Length
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                                                                                                                 Gaps
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RESULT 9
AAR99259
ID AAR99259
XX AAR9
XX AAR9
AC AAR9
XX CD40
XX CD40
KW CD40
KW CD40
KW CD40
KW CD40
KW CD40
XX HOMO
XX HOMO
XX HOMO
XX HOMO
XX HOMO
XX I LOMB
FT DOMB
FT NOB6
XX WO96
XX WO96
XX NOB6
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A novel human CD40 receptor binding protein (CD40bp) (AAR99259) has the ability to bind the cytoplasmic region of the CD40 receptor. Its amino acid sequence was deduced from a cDNA clone (AAT35251)
                                                                                                                                                                                                                                        N-PSDB; AAT35251.
                                                                                                                                                                                                                                                                   WPI; 1996-433838/43
                                                                                                                                                                                                                                                                                                                         Dixit VM;
                                                                                                           Example 5; Page 41-43; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                         (UNMI ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                            13-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9628568-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibody; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CD40 binding protein; CD40bp; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Full-length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR99259 standard;
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                                                                                                                                                   isolated CD40 receptor binding protein - used to develop prods. use as immunosuppressive drugs and to treat immune disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TVLENGTYIKDDTIFIKVIVDTSDLPDP 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TVLENGTY IKDDT IF IKVIVDTSDLPDP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VTELESVDKSAGQAARNTGLLESQLSRHDQTLSVHDIRLADMDLRFQVLETASYNGVLIW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          \tt VTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLGFQVLETASYNGVLIW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CD40 binding protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                         MICHIGAN
                                                                                                                                                                                                                                                                                                                                                                                                                            95US-0404832
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95WO-US06623
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/label= RING_finger_domain
266..376
/label= Coiled-coil_domain
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CD40 receptor associated factor 1; CRAF1-a; TRAF-3; p55; human; CD40 mediated intracellular signalling; organ rejection; allergy; hay fever; autoimmune disease; systemic lupus erythematosus; rheumatoid arthritis; myasthenia gravis; Graves' disease; diopathic thrombocytopaenia purpura; haemolytic anaemia; diabetes mellitus; psoriasis; hyper immunoglobulin E syndrome; apoptosis; Rieter's syndrome; spondyloarthritis; Lyme disease; HIV; syphilis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis;
                                                                                                                                                                                                                                                                      Human CRAF1 isoform p55del9
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Pred. No. 6.3e-229;
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This polypeptide comprises a CRAFI (TRAF-3) isoform designated CC p55del9. It is encoded by exons 4-8 and 10-13 of the human CRAF CC gene (see AAM790123) and arises by alternative splicing of the sequence for CRAF1-a (see AAW27431), a signalling protein that CC interacts with the cytoplasmic tail of B cell surface molecule CD40 and which mediates a variety of T-dependent effects on B cell cactivation and differentiation. A higher mol.wt. CRAF1 designated CC CRAF1b (see AAW27428), has also been identified, as well as isoforms CC post of the see AAW27429), p15 (see AAW27430) and variants of CRAF-1a and CC CRAF-1b (see AAW27432-37) that comprise different combinations of 5 (see AAW27432-37) that comprise different combinations of 5 (companies). CRAF1 peptides, comprising from 0-4 zinc finger CC domains, and nucleic acids encoding them, can be used to inhibit CC do ligand activation of cells that express CD40 on their surface, particularly by introducing a nucleic acid molecule into the cells, and used to treat conditions characterised by an aberrant or companies of the companies o
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21-MAR-1996;
01-MAY-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protein comprising CRAFI-b domain capable of inhibiting CD40 mediated cell activation - useful to treat conditions characterised by aberrant or unwanted level of CD40 mediated intracellular
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associated with Epstein-Barr virus
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asbestosis; siliconosis; Farmer's lung; hepatitis; cirrhosis;
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KW CD40 receptor associated factor 1; CRAF1-b; TRAF-3; p70; human;
KW CD40 mediated intracellular signalling; organ rejection; allergy;
KW hay fever; autoimmune disease; systemic lupus erythematosus;
KW rheumatoid arthritis; myasthenia gravis; Graves' disease;
KW idiopathic thrombocytopaenia purpura; haemolytic anaemia;
KW diabetes mellitus; psoriasis; hyper immunoglobulin E syndrome;
KW apoptosis; Rieter's syndrome; spondyloarthritis; Lyndrome; promonitis;
KW syphilis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis;
KW syphilis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis;
KW syphilis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis;
KW sphilis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis;
KW shestosis; siliconosis; Farmer's lung; hepatitis; cirrhosis;
KW atherosclerosis; multiple sclerosis; glomerulonephritis;
KW atherosclerosis; elomerulopathy; kidney disease; nephropathy;
KW endocarditis; leprosy; malaria; Goodpasture's disease;
KW Henoch-Schoenlein purpura; polyarteritis; multiple myeloma;
KW Wegener's granulomatosis; cryoimmunoglobulinaemia;
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Best Local
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Pred. No. 1.1e-218;
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Best Local Similarity

91.7%;

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18-SEP-1996;
21-MAR-1996;
01-MAY-1996;
                                        treat conditions characterised by an aberrant or unwanted level of CD40 mediated intracellular signalling, such as organ rejection, or a CD40 dependent immune response in a subject receiving gene therapy. The condition may be an allergic response or an autoimmune response, or may be dependent on CD40 ligand-induced activation of epithelial cells, an inflammatory kidney disease, a smooth muscle cell-dependent disease, or a condition associated
                                                                                                                                                 This polypeptide comprises a CRAF1 (TRAF-3) protein designated p70del9 that is encoded by exons 1-2, 4-8 and 10-13 of the human CRAF gene (see AAT90123). Different isoforms (AAW27428-37) of CRAF1 have been identified that arise from alternative splicing. CRAF1 peptides comprising from 0-4 zinc finger domains, and nucleic acids encoding them, can be used to inhibit CD40 ligand activation of cells that express CD40 on their surface, particularly by introducing the nucleic acid molecule into the cells, and used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Waldenstroem's macroglobulinaemia; amyloidosis; Sjogren's sydrome; AIDS; oesophageal dysmotility; inflammatory bowel disease; bladder disease; Egstein-Barr virus; mononucleosis; B cell tumour; Burkitt's lymphoma; nasopharyngeal carcinoma; pneumonia;
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Sequence
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                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; AAT90123
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Cleary AM, Frank DM,
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                             with Epstein-Barr virus.
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/note= "Claim 1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                          Lederman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "putative SH3 binding motif"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "putative SH3 binding motif"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "zinc finger 2 (zinc binding to Cys-270, Cys-275, His-287 and Cys-292 "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "zinc finger 1 (2n binding to Cys-246, His-258 and Cys-263)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "zinc finger 3 (Zn binding to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "putative SH3 binding motif"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cys-302, His-314 and Cys-319"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       _Cys-299
                                                                                                                                                                                                                                                                                                                                                    characterised
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Query Match

91.0%;

Score 2726;

DB 18;

Length 665;

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RESULT 12
AAR98833
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                                                                                                                                                                                                                                                   CD40 associated protein (CAP)-1.
                                                 Domain
                                                                                                 Domain
                                                                                                                          Domain
                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                             23-MAR-1998
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             WO9616665-A1
                                                                        Domain
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                                                                                                                                                                                                               associated protein; CAP; agonist; antagonist; gene therapy; proliferation; treatment; cancer; autoimmune disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     EYDALLPWPFKQKYTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ 539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VTELESVDKSAGQAARNTGLLESQLSRHDQTLSVHDIRLADMDLRFQVLETASYNGVLIW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EIERQKEMLRUNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        --GTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSE----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKYREATCSHC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CSPKQTECGHRFCESCMAALLSSSSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
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                                                                                                                                                                                                                                                                            (first entry)
                                                                                                 /note= "2
148..170
                                                                          /note=
177..1
                                                                                                                                                 Location/Qualifiers 53..91
                                                /note= "Zinc finger domain
384..540
                                                                                                                          /note= "RING finger domain"
117..141
                                                                                                                                                                                                                                                                                                                               Protein; 543 AA
                                                                          ..197
                                                                                      "Zinc
                                                                                                             "Zinc finger domain 1"
                                     "TRAF domain"
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                                                                                    finger
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This is a CD40 associated protein (CAP)-1. This CAP is a protein that specifically binds to CD40, a cell surface receptor involved in apoptosis. Agonists and antagonists of CAP can increase or decrease the level of CAP expression in a cell and can thereby modulate the function of the cell. Such compounds can be used to treat cancer, autoimmune diseases like asthma, hay fever, rheumatoid arthritis and immunodeficiency diseases and neurodegeneration. Antibodies that bind specifically to CAP can be used to assay CAP, to detect pathologically altered levels. The encoding nucleic acid can be used to identify related genes and to express CAP for gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New CD40 associated protein, agonists and antagonists - used modulate cell proliferation, immune response, apoptosis etc., for treating cancer or autoimmune disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 MESSKKMDAAGTLQPNPPLKLQPDRGAGS-VLVPEQGGYKEKFVKTVEDKYKCEKCRLVL 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHC 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CNPKQTECGHRFCESCMAALLSSSSPKCTACQESIIKDKVFKDNCCKREILALQVYCRNE 119
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DB; AAT30773.
TVLENGTYIKDDTIFIKVIVDTSDLPDP 567
                                                                                                                    KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
                                                                                                                                                                                      VTELESVDKSAGQAARNTGLLESQLSRHDQTLSVHDIRLADMDLRFQVLETASYNGVLIW 419
                                                                                                                                                                                                                                                           EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR 359
                                                                                                                                                                                                                                                                                                       --GTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI 275
                                  EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ
                                                                EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ 539
                                                                                                  KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
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                                                                                                                                                                     VTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLRFQVLETASYNGVLIW
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91.4%;
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RESULT 13
AAW27436
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KW CD40 mediated intracellular signalling; organ rejection; allergy;
KW chay fever; autoimmune disease; systemic lupus erythematosus;
KW rheumatoid arthritis; myasthenia gravis; Graves' disease;
KW idiopathic thrombocytopaenia purpura; haemolytic anaemia;
KW apoptosis; Rieter's syndrome; spondyloarthritis; Lyme disease; HIV;
KW apoptosis; Rieter's syndrome; spondyloarthritis; Lyme disease; HIV;
KW syphilis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis;
KW ppeumoconiosis; adult respiratory distress syndrome; pneumonitis;
KW pneumoconiosis; adult respiratory distress syndrome; pneumonitis;
KW preschesis; siliconosis; Farmer's lung; hepatitis; cirrhosis;
KW atherosclerosis; multiple sclerosis; glomerulonephritis;
KW glomerulosclerosis; glomerulopathy; kidney disease; nephropathy;
KW Henoch-Scheenlein purpura; polyarteritis; multiple myeloma;
KW Wegener's granulomatosis; cryolmmunoglobulinaemia;
KW Waldenstroem's macroglobulinaemia; amyloidosis; Slogren's sydrome;
KW Waldenstroem's macroglobulinaemia; amyloidosis; Slogren's sydrome;
                                                                                                                               18-SEP-1996;
21-MAR-1996;
01-MAY-1996;
01-MAY-1996;
Protein comprising CRAF1-b domain capable of inhibiting CD40
                            WPI; 1997-479907/44.
N-PSDB; AAT90123.
                                                                       Cleary
                                                                                                                                                                                                                                                                                                       Binding-site
                                                                                                                                                                                                                                                                                                                                                               Binding-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AIDS; oesophageal dysmotility; inflammatory bowel disease; bladder disease; Epstein-Barr virus; mononucleosis; B cell tumour; Burkitt's lymphoma; nasopharyngeal carcinoma; pneumonia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW27436
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                                                                                                 (UYCO ) UNIV COLUMBIA NEW YORK.
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                                                                                                                                                                                                                                                                                                                                                                                                          Region
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                                                                                                                                                                                                                                                              WO9734473-A1
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                                                                       Frank DM,
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                                                                                                                               96US-0013820.
96US-0016626.
96US-0016659.
                                                                                                                                                                         96US-0026584
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                                                                                                                                                                                                                                                                                                       /note= "putative SH3 binding motif"
103..110
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Cys-275, His-287 and Cys-292
                                                                                                                                                                                                                                                                                                                                                  "putative SH3 binding
                                                                                                                                                                                                                                                                                                                                                                                                                        "zinc finger 1 (Zn binding to Cys-239, Cys-246, His-258 and Cys-263)"
                                                                                                                                                                                                                                                                                           "putative SH3 binding motif"
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"Claim 1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This polypeptide comprises a CRAFI (TRAF-3) protein designated 20 p70del8,9 that is encoded by exons 1-2, 4-7 and 10-13 of the human 21 CRAF gene (see AAT90123). Different isoforms (AAW27428-37) of CRAFI 22 CRAF gene (see AAT90123). Different isoforms (particular) of CRAFI 23 cross periods comprising from 0-4 zinc finger domains, and nucleic acids 25 centred to the m, can be used to inhibit CD40 ligand activation of 26 cells that express CD40 on their surface, particularly by 27 cells that express CD40 on their surface, particularly by 28 cells that express CD40 on their surface, particularly by 29 cells that express CD40 on their surface, particularly by 29 cells that express CD40 on their surface, particularly by 29 cells that express conditions characterised by an aberrant or unwanted level of 20 cells that express a signalling, such as organ rejection, or 20 cells condition may be an allergic response or an 20 cell-dependent immune response or an allergic response or an 20 cells condition of epithelial cells, an inflammatory kidney disease, a 20 cells cell-dependent disease, or a condition associated
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 489
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TVLENGTYIKDDTIFIKVIVDTSDLPDP
                                                                                                                                                                         KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
                                                                                                                                                                                                                                               VTELESVDKSAGQAARNTGLLESQLSRHDQTLSVHDIRLADMDLRFQVLETASYNGVLIW
                                                                                                                                                                                                                                                                                                                            EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR
                                                                                                                                                                                                                                                                                                                                                                                                     FQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MESSKKMDSPGALQTNPPLKLHTDRSAGTPVFVPEQGGYKEKFVKTVEDKYKCEKCHLVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MESSKKMDAAGTLQPNPPLKLQPDRGAGS-VLVPEQGGYKEKFVKTVEDKYKCEKCRLVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Epstein-Barr virus.
                                   TVLENGTYIKDDTIFIKVIVDTSDLPDP 567
                                                                        EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ
                                                                                                              EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNTASGCPVFVAQ
                                                                                                                                                   KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFYIMRG
                                                                                                                                                                                                                              VTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLRFQVLETASYNGVLIW
                                                                                                                                                                                                                                                                                                      EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR
                                                                                                                                                                                                                                                                                                                                                                                                                                                            KSQVPMIAL----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KSQVPMIKLQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CSPKQTECGHRFCESCMAALLSSSSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CNPKQTECGHRFCESCMAALLSSSSPKCTACQESIIKDKVFKDNCCKREILALQVYCRNE
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87.0%;
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Pred. No. 9.7e-205;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
 516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 18; Length 516;
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RESULT 14
AAW27437
ID AAW27

AAW27437

standard;

Protein;

638

This polypeptide comprises a CRAF1 (TRAF-3) protein designated

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18-SEP-1996;
21-MAR-1996;
01-MAY-1996;
01-MAY-1996;
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                                                               Protein comprising CRAF1-b domain capable of inhibiting CD40 mediated cell activation - useful to treat conditions charact by aberrant or unwanted level of CD40 mediated intracellular
                                                                                                                                                               Cleary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AIDS; oesophageal dysmotility; inflammatory bowel disease; bladder disease; Epstein-Barr virus; mononucleosis; B cell Burkitt's lymphoma; nasopharyngeal carcinoma; pneumonia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CD40 receptor associated factor 1; CRAF1-b; TRAF-3; p70; human; CD40 mediated intracellular signalling; organ rejection; allergy; hay fever; autoimmune disease; systemic lupus erythematosus; rheumatoid arthritis; myasthenia gravis; Graves' disease; idiopathic thrombocytopaenia purpura; haemolytic anaemia; diabetes mellitus; psoriasis; hyper immunoglobulin E syndrome;
                        Example 1; Fig 1A-O;
                                                                                                                       WPI; 1997-479907/44.
N-PSDB; AAT90123.
                                                                                                                                                                                                                                                                                         21-MAR-1997;
                                                                                                                                                                                                                                                                                                                    25-SEP-1997
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                                                                                                                                                                                                                                                                                                                                               WO9734473-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Domain
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                                                                                                                                                               Frank DM,
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96US-0013820.
96US-0016626.
96US-0016659.
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                                                                                                                                                                                                                                                                                                                                                                         103..1
/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /label= CRAF-b_domain
/note= "Claim 1"
239..263
                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "putative SH3 binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
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                      158pp;
                                                                                                                                                                Lederman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "zinc finger 1 (Zn binding to Cys-239, Cys-246, His-258 and Cys-263)"
                                                                                                                                                                                                                                                                                                                                                                         "putative
                                                                                                                                                                                                                                                                                                                                                                                                                                                         "zinc finger 2 (zinc binding Cys-275, His-287 and Cys-292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cys-246,
                        English
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                                                                                                                                                                                                                                                                                                                                                                           SH3 binding
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                                                                  ditions characterised intracellular
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RESULT 15
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                              Human CRAF1
                                                                                                                                                           AAW27434
                                                                                           27-MAR-1998
                                                                                                                                                                                                                     AAW27434 standard; Protein; 512
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                                                                                                                                                                                                                                                                                                                                                                             TVLENGTYIKDDTIFIKVIVDTSDLPDP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLRFQVLETASYNGVLIW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEADSMKSSVESLQNR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKYREATCSHC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CSPKQTECGHRFCESCMAALLSSSSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KSQVPMIKLQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CNPKQTECGHRFCESCMAALLSSSSPKCTACQESIIKDKVFKDNCCKREILALQVYCRNE 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -QGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Epstein-Barr virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                638 AA;
                           isoform p55del9,10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 18; Length 638;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          610
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KW CD40 receptor associated factor 1; CRAF1-a; TRAF-3; p55; human; KW CD40 mediated intracellular signalling; organ rejection; allergy; KW chemicatoid arthritis; myasthenia gravis; Graves' disease; KW rheumatoid arthritis; myasthenia gravis; Graves' disease; KW idiopathic thrombocytopaenia purpura; haemolytic anaemia; Midopathic thrombocytopaenia purpura; haemolytic anaemia; KW diabetes mellitus; psoriasis; hyper immunoglobulin E syndrome; KW apoptosis; Rieter's syndrome; spondyloarthritis; Lyme disease; HIV; KW syphilis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis; KW syphilis; tuberculosis; arthritis; scleroderma; pulmonary fibrosis; KW appetosis; siliconosis; Farmer's lung; hepatitis; cirrhosis; KW abbestosis; siliconosis; Farmer's lung; hepatitis; cirrhosis; KW atherosclerosis; multiple sclerosis; glomerulonephritis; Malbestosis; leprosy; malaria; Goodpasture's disease; Menoch-Schoenlein purpura; polyarteritis; multiple myeloma; KW wegener's macroglobulinaemia; amyloldosis, Sjogren's sydrome; KW Waldenstroem's macroglobulinaemia; amyloldosis; Sjogren's sydrome; KW MiDS; oesophageal dysmotility; inflammatory bowel disease; Bpstein Barr virus; mononucleosis; B cell tumour; KW bladder disease; Epstein Barr virus; mononucleosis; B cell tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-SEP-1996;
21-MAR-1996;
01-MAY-1996;
01-MAY-1996;
                                                                                                                                                                                                                                                                              Protein comprising CRAF1-b domain capable of inhibiting CD40 mediated cell activation - useful to treat conditions charac by aberrant or unwanted level of CD40 mediated intracellular
                                                                                                                                                                                                                                                                                                                                                                                             Cleary AM,
                                                                                                                                                                                                                                                                                                                                                N-PSDB; AAT90123.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-SEP-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                           (UYCO ) UNIV COLUMBIA NEW YORK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                               1997-479907/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy; diagnosis
                                                                                                                                                                                                                                                                                                                                                                                               Frank DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         96US-0013820.
96US-0016626.
96US-0016659.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96US-0026584.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97WO-US05076
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "zinc finger 2 (zinc binding to Cys-148, Cys-153, His-165 and Cys-170"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                             Lederman S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "zinc finger 3 (Zn binding to Cys-177 Cys-180, His-192 and Cys-197"
                                                                                                                                                                                                                                                                                               characterised
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Example 1; Fig 1D-O; 158pp; English.

This polypeptide comprises a CRAFI (TRAF-3) isoform designated pp55del9,10. It is encoded by exons 4-8 and 11-13 of the human CRAF gene (see AAT90123) and arises by alternative splicing of the sequence for CRAF1-a (see AAW27431), a signalling protein that interacts with the cytoplasmic tail of B cell surface molecule CD40 and which mediates a variety of T-dependent effects on B cell activation and differentiation. A higher mol.wt. CRAF1, designated CRAF1b (see AAW27428), has also been identified, as well as isoforms p5 (see AAW27429), p15 (see AAW27430) and variants of CRAF-la and CRAF-b (see AAW2742-37) that comprise different combinations of 5 zinc fingers. CRAF1 peptides, comprising from 0-4 zinc finger domains, and nucleic acids encoding them, can be used to inhibit counter the combination of cells that express CD40 in their surface, particularly by introducing a nucleic acid molecule into the cells

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Best Local S
Matches 490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and used to treat conditions characterised by an aberrant or unwanted level of CD40 mediated intracellular signalling, such as organ rejection, or a CD40 dependent immune response in a subject receiving gene therapy. The condition may be an allergic response or an autoimmune response, or may be dependent on CD40 ligandinduced activation of epithelial cells, an inflammatory kidney disease, a smooth muscle cell-dependent disease, or a condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             associated with Epstein-Barr virus.
             540 TVLENGTYIKDDTIFIKVIVDTSDLPDP 567
                                                        425 EYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQ 484
                                                                                   480
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 MESSKKMDAAGTLQPNPPLKLQPDRGAGS-VLVPEQGGYKEKFVKTVEDKYKCEKCRLVL 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                        KSQVPMIKLQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCV 239
                                                                      EYDALLPWPFKOKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKFTGEMNIASGCPVFVAQ 539
                                                                                                              KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
                                                                                                                              KIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRG
                                                                                                                                                                                                                                                                                                                 FQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI 299
                                                                                                                                                                                                                                                                                                                                          KSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSE-----
                                                                                                                                                                                                                                                                                                                                                                                                    SRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKYREATCSHC 180
                                                                                                                                                                                                                                                                                                                                                                                                                  GRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHC 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TVLENGTYIKDDTIFIKVIVDTSDLPDP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        490;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 512 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      85.3%;
86.3%;
                                                                                                                                                                                                                                                                                      -----VSLLQNESVEKNKSIQSLHNQICSFEI 244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 2552.5; DB Pred. No. 3e-204; 7; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 18; Length 512;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14; Indels
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                                                                                                                 424
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Search completed: December 19, 2002, 15:00:24 Job time : 40 secs